

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/093,972

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2 Wrapped Aminos The amino acid number/text at the end of each line "wrapped " down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3 ✓ Incorrect Line Length The rules require that a line not exceed 72 characters in length. This includes spaces.
- 4 Misaligned Amino Acid The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs
Numbering between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
- 5 Non-ASCII This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6 Variable Length Sequence(s) contain n's or Xaa's which represented more than one residue.
As per the rules, each n or Xaa can only represent a single residue.
Please present the maximum number of each residue having variable length and
indicate in the (ix) feature section that some may be missing.
- 7 PatentIn ver. 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid
sequence(s) . Normally, PatentIn would automatically generate this section from the
previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section
to the subsequent amino acid sequence.
- 8 Skipped Sequences Sequence(s) missing. If intentional, please use the following format for each skipped sequence:
(OLD RULES) (2) INFORMATION FOR SEQ ID NO:X:
 (i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:
 This sequence is intentionally skipped

Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9 Skipped Sequences Sequence(s) missing. If intentional, please use the following format for each skipped sequence.
(NEW RULES) <210> sequence id number
 <400> sequence id number
 000
- 10 Use of n's or Xaa's Use of n's and/or Xaa's have been detected in the Sequence Listing.
(NEW RULES) Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 11 Use of <213>Organism Sequence(s) are missing this mandatory field or its response.
(NEW RULES)
- 12 Use of <220>Feature Sequence(s) are missing the <220>Feature and associated headings.
(NEW RULES) Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
- 13 PatentIn ver. 2.0 "bug" Please do not use "C py to Disk" function of PatentIn version 2.0. This causes a corrupted
file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).
Instead, please use "File Manager" or any other means to copy file to floppy disk.
AKS-Biotechnology Systems Branch- 5/15/99

EPFS

1635

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/093,972

DATE: 12/16/1999
TIME: 03:36:59

INPUT SET: S34241.raw

This Raw Listing contains the General Information Section and those Sequences containing ERRORS.

Does Not Comply
Corrected Diskette Needed

SEQUENCE LISTING

1
2
3 (1) General Information:
4 (i) APPLICANT: Nyce, Jonathan W.
5 (ii) TITLE OF INVENTION: COMPOSITION, FORMULATIONS & METHOD FOR PREVENTION & T
6 (iii) NUMBER OF SEQUENCES: 996
7 (iv) CORRESPONDENCE ADDRESS:
8 (A) ADDRESSEE: ARTER & HADDEN
9 (B) STREET: 725 South Figueroa Street
10 (C) CITY: Los Angeles
11 (D) STATE: California
12 (E) COUNTRY: USA
13 (F) ZIP: 90017
14 (v) COMPUTER READABLE FORM:
15 (A) MEDIUM TYPE: Floppy disk
16 (B) COMPUTER: IBM PC compatible
17 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
18 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
19 (vi) CURRENT APPLICATION DATA:
20 (A) APPLICATION NUMBER: US 09/093,972 OK
21 (B) FILING DATE: 9-JUNE-1998
22 (C) CLASSIFICATION:
23 (viii) ATTORNEY/AGENT INFORMATION:
24 (A) NAME: Amzel, Viviana
25 (B) REGISTRATION NUMBER: 30,930
26 (C) REFERENCE/DOCKET NUMBER: EPI-072 (73999\95804)
27 (ix) TELECOMMUNICATION INFORMATION:
28 (A) TELEPHONE: 213-430-3520
29 (B) TELEFAX: 213-617-9255
30 (C) TELEX:
31

all text
must be
visible
see item 3
on Encl
summary
sheet

ERRORED SEQUENCES FOLLOW:

9566 (2) INFORMATION FOR SEQ ID NO:953:
9567 (i) SEQUENCE CHARACTERISTICS:
--> 9568 (A) LENGTH: 23 base pairs
9569 (B) TYPE: nucleic acid
9570 (C) STRANDEDNESS: single
9571 (D) TOPOLOGY: linear
9572 (ii) MOLECULE TYPE: DNA (genomic)
9573 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:953:

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/093,972DATE: 12/16/1999
TIME: 03:36:59

INPUT SET: S34241.raw

23 ← move over from far right

9574 TTT TCC TTC CTT TGT CTC TCT TC
9575

--> 9576 (2) INFORMATION FOR SEQ ID NO:954:
9577 (i) SEQUENCE CHARACTERISTICS:
9578 (A) LENGTH: 15 base pairs
9579 (B) TYPE: nucleic acid
9580 (C) STRANDEDNESS: single
9581 (D) TOPOLOGY: linear
9582 (ii) MOLECULE TYPE: DNA (genomic)
9583 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:954:
9584 GCT CCC GGC TGC CTG
9585

same env

--> 9586 (2) INFORMATION FOR SEQ ID NO:955:
9587 (i) SEQUENCE CHARACTERISTICS:
9588 (A) LENGTH: 29 base pairs
9589 (B) TYPE: nucleic acid
9590 (C) STRANDEDNESS: single
9591 (D) TOPOLOGY: linear
9592 (ii) MOLECULE TYPE: DNA (genomic)
9593 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:955:
9594 CTC GGC CGT GCG GCT CTG TCG CTC CCG GT
9595

same

--> 9596 (2) INFORMATION FOR SEQ ID NO:956:
9597 (i) SEQUENCE CHARACTERISTICS:
9598 (A) LENGTH: 20 base pairs
9599 (B) TYPE: nucleic acid
9600 (C) STRANDEDNESS: single
9601 (D) TOPOLOGY: linear
9602 (ii) MOLECULE TYPE: DNA (genomic)
9603 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:956:
9604 CCG CCG CCC TCC GGG GGG TC
9605

same

--> 9606 (2) INFORMATION FOR SEQ ID NO:957:
9607 (i) SEQUENCE CHARACTERISTICS:
9608 (A) LENGTH: 18 base pairs
9609 (B) TYPE: nucleic acid
9610 (C) STRANDEDNESS: single
9611 (D) TOPOLOGY: linear
9612 (ii) MOLECULE TYPE: DNA (genomic)
9613 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:957:
9614 TGC TGC CGT TGG CTG CCC
9615

same

--> 9616 (2) INFORMATION FOR SEQ ID NO:958:
9617 (i) SEQUENCE CHARACTERISTICS:
9618 (A) LENGTH: 17 base pairs
9619 (B) TYPE: nucleic acid
9620 (C) STRANDEDNESS: single
9621 (D) TOPOLOGY: linear

RAW SEQUENCE LISTING
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9622 (ii) MOLECULE TYPE: DNA (genomic)
9623 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:958:
9624 CTT CTG CGG GTC GCC GG
9625

same

--> 9626 (2) INFORMATION FOR SEQ ID NO:959:
9627 (i) SEQUENCE CHARACTERISTICS:
9628 (A) LENGTH: 15 base pairs
9629 (B) TYPE: nucleic acid
9630 (C) STRANDEDNESS: single
9631 (D) TOPOLOGY: linear
9632 (ii) MOLECULE TYPE: DNA (genomic)
9633 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:959:
9634 TGC TGG GCT TGT GGC
9635

same

--> 9636 (2) INFORMATION FOR SEQ ID NO:960:
9637 (i) SEQUENCE CHARACTERISTICS:
9638 (A) LENGTH: 15 base pairs
9639 (B) TYPE: nucleic acid
9640 (C) STRANDEDNESS: single
9641 (D) TOPOLOGY: linear
9642 (ii) MOLECULE TYPE: DNA (genomic)
9643 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:960:
9644 GGC CTC TCT TCT GGG
9645

same

--> 9646 (2) INFORMATION FOR SEQ ID NO:961:
9647 (i) SEQUENCE CHARACTERISTICS:
9648 (A) LENGTH: 14 base pairs
9649 (B) TYPE: nucleic acid
9650 (C) STRANDEDNESS: single
9651 (D) TOPOLOGY: linear
9652 (ii) MOLECULE TYPE: DNA (genomic)
9653 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:961:
9654 CCT GGT CCC TCC GT
9655

same

--> 9656 (2) INFORMATION FOR SEQ ID NO:962:
9657 (i) SEQUENCE CHARACTERISTICS:
9658 (A) LENGTH: 14 base pairs
9659 (B) TYPE: nucleic acid
9660 (C) STRANDEDNESS: single
9661 (D) TOPOLOGY: linear
9662 (ii) MOLECULE TYPE: DNA (genomic)
9663 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:962:
9664 GGT GGC TCC TCT GC
9665

same

--> 9666 (2) INFORMATION FOR SEQ ID NO:963:
9667 (i) SEQUENCE CHARACTERISTICS:
9668 (A) LENGTH: 18 base pairs
9669 (B) TYPE: nucleic acid

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/093,972DATE: 12/16/1999
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INPUT SET: S34241.raw

9670 (C) STRANDEDNESS: single
9671 (D) TOPOLOGY: linear
9672 (ii) MOLECULE TYPE: DNA (genomic)
9673 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:963:
9674 GCT TGG TCC TGG GGC TGC *same*
9675

--> 9676 (2) INFORMATION FOR SEQ ID NO:964:
9677 (i) SEQUENCE CHARACTERISTICS:
9678 (A) LENGTH: 15 base pairs
9679 (B) TYPE: nucleic acid
9680 (C) STRANDEDNESS: single
9681 (D) TOPOLOGY: linear
9682 (ii) MOLECULE TYPE: DNA (genomic)
9683 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:964:
9684 TGC TCT CCT CTC CTT *same*
9685

--> 9686 (2) INFORMATION FOR SEQ ID NO:965:
9687 (i) SEQUENCE CHARACTERISTICS:
9688 (A) LENGTH: 21 base pairs
9689 (B) TYPE: nucleic acid
9690 (C) STRANDEDNESS: single
9691 (D) TOPOLOGY: linear
9692 (ii) MOLECULE TYPE: DNA (genomic)
9693 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:965:
9694 TGC TTT TCT TTT CTG GGC CTC *same*
9695

--> 9696 (2) INFORMATION FOR SEQ ID NO:966:
9697 (i) SEQUENCE CHARACTERISTICS:
9698 (A) LENGTH: ~~18~~ base pairs → 19 shown
9699 (B) TYPE: nucleic acid
9700 (C) STRANDEDNESS: single
9701 (D) TOPOLOGY: linear
9702 (ii) MOLECULE TYPE: DNA (genomic)
9703 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:966:
9704 TGT GGT CTG TTT TTT TCT G-3= delete 19C
9705

--> 9706 (2) INFORMATION FOR SEQ ID NO:967:
9707 (i) SEQUENCE CHARACTERISTICS:
9708 (A) LENGTH: 20 base pairs
9709 (B) TYPE: nucleic acid
9710 (C) STRANDEDNESS: single
9711 (D) TOPOLOGY: linear
9712 (ii) MOLECULE TYPE: DNA (genomic)
9713 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:967:
9714 GCC CTG CTG GGG CGC TCT CC-3= *same error*
9715

9716 (2) INFORMATION FOR SEQ ID NO:968:
9717 (i) SEQUENCE CHARACTERISTICS:

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/093,972DATE: 12/16/1999
TIME: 03:37:00

INPUT SET: S34241.raw

--> 9718 (A) LENGTH: 18 base pairs
9719 (B) TYPE: nucleic acid
9720 (C) STRANDEDNESS: single
9721 (D) TOPOLOGY: linear
9722 (ii) MOLECULE TYPE: DNA (genomic)
9723 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:968:
9724 GCC GCC CGC CTG GCT CCC(3)
9725

same error

9726 (2) INFORMATION FOR SEQ ID NO:969:
9727 (i) SEQUENCE CHARACTERISTICS:
--> 9728 (A) LENGTH: 21 base pairs
9729 (B) TYPE: nucleic acid
9730 (C) STRANDEDNESS: single
9731 (D) TOPOLOGY: linear
9732 (ii) MOLECULE TYPE: DNA (genomic)
9733 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:969:
9734 GGB GCC CBT GBT GGG CBT GCC
9735

21 ← move

9736 (2) INFORMATION FOR SEQ ID NO:970:
9737 (i) SEQUENCE CHARACTERISTICS:
--> 9738 (A) LENGTH: 24 base pairs
9739 (B) TYPE: nucleic acid
9740 (C) STRANDEDNESS: single
9741 (D) TOPOLOGY: linear
9742 (ii) MOLECULE TYPE: DNA (genomic)
9743 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:970:
9744 GTG GTT CTT GCC CTC CTT TGG CTG
9745

same error

9746 (2) INFORMATION FOR SEQ ID NO:971:
9747 (i) SEQUENCE CHARACTERISTICS:
--> 9748 (A) LENGTH: 18 base pairs
9749 (B) TYPE: nucleic acid
9750 (C) STRANDEDNESS: single
9751 (D) TOPOLOGY: linear
9752 (ii) MOLECULE TYPE: DNA (genomic)
9753 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:971:
9754 CCG TGC CCG CTC CCC GGC
9755

same

9756 (2) INFORMATION FOR SEQ ID NO:972:
9757 (i) SEQUENCE CHARACTERISTICS:
--> 9758 (A) LENGTH: 20 base pairs
9759 (B) TYPE: nucleic acid
9760 (C) STRANDEDNESS: single
9761 (D) TOPOLOGY: linear
9762 (ii) MOLECULE TYPE: DNA (genomic)
9763 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:972:
9764 CTC CTG GCG GGT GGC CGT TG
9765

same

9766 (2) INFORMATION FOR SEQ ID NO:973:

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/093,972

DATE: 12/16/1999
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INPUT SET: S34241.raw

--> 9767 (i) SEQUENCE CHARACTERISTICS:
9768 (A) LENGTH: 18 base pairs
9769 (B) TYPE: nucleic acid
9770 (C) STRANDEDNESS: single
9771

ignore blank section

(D) TOPOLOGY: linear

9772 (ii) MOLECULE TYPE: DNA (genomic)
9773 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:973:
9774 GGC CCG TGT TCC CCT GGG
9775

same error

--> 9776 (2) INFORMATION FOR SEQ ID NO:974:
9777 (i) SEQUENCE CHARACTERISTICS:
9778 (A) LENGTH: 20 base pairs
9779 (B) TYPE: nucleic acid
9780 (C) STRANDEDNESS: single
9781 (D) TOPOLOGY: linear
9782 (ii) MOLECULE TYPE: DNA (genomic)
9783 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:974:
9784 GCC TGG GGC TCC CTT CTC TC
9785

same

--> 9786 (2) INFORMATION FOR SEQ ID NO:975:
9787 (i) SEQUENCE CHARACTERISTICS:
9788 (A) LENGTH: 19 base pairs
9789 (B) TYPE: nucleic acid
9790 (C) STRANDEDNESS: single
9791 (D) TOPOLOGY: linear
9792 (ii) MOLECULE TYPE: DNA (genomic)
9793 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:975:
9794 GCC CTT CTT GCT GGG CCT C
9795

same

--> 9826 (2) INFORMATION FOR SEQ ID NO:979:
9827 (i) SEQUENCE CHARACTERISTICS:
9828 (A) LENGTH: 29 base pairs
9829 (B) TYPE: nucleic acid
9830 (C) STRANDEDNESS: single
9831 (D) TOPOLOGY: linear
9832 (ii) MOLECULE TYPE: DNA (genomic)
9833 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:979:
9834 GGC GCC GTG CCG CGT CTT GGT GGC GGC GG
9835

same

--> 9836 (2) INFORMATION FOR SEQ ID NO:980:
9837 (i) SEQUENCE CHARACTERISTICS:
9838 (A) LENGTH: 30 base pairs
9839 (B) TYPE: nucleic acid
9840 (C) STRANDEDNESS: single
9841 (D) TOPOLOGY: linear
9842 (ii) MOLECULE TYPE: DNA (genomic)
9843 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:980:
9844 GTT CGC GCC CGC GCG GGG CCC CTC CGG TCC
9845

same

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/093,972DATE: 12/16/1999
TIME: 03:37:01

INPUT SET: S34241.raw

9856 (2) INFORMATION FOR SEQ ID NO:982:
9857 (i) SEQUENCE CHARACTERISTICS:
--> 9858 (A) LENGTH: 22 base pairs
9859 (B) TYPE: nucleic acid
9860 (C) STRANDEDNESS: single
9861 (D) TOPOLOGY: linear
9862 (ii) MOLECULE TYPE: DNA (genomic)
9863 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:982:
9864 CGG GTC GGG GCC CCC CGC GGC C
9865

same

9866 (2) INFORMATION FOR SEQ ID NO:983:
9867 (i) SEQUENCE CHARACTERISTICS:
--> 9868 (A) LENGTH: 29 base pairs
9869 (B) TYPE: nucleic acid
9870 (C) STRANDEDNESS: single
9871 (D) TOPOLOGY: linear
9872 (ii) MOLECULE TYPE: DNA (genomic)
9873 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:983:
9874 GCC TCG GGG CTG GGG CGC TGG TGG CCG GG
9875

same

9876 (2) INFORMATION FOR SEQ ID NO:984:
9877 (i) SEQUENCE CHARACTERISTICS:
--> 9878 (A) LENGTH: 24 base pairs
9879 (B) TYPE: nucleic acid
9880 (C) STRANDEDNESS: single
9881 (D) TOPOLOGY: linear
9882 (ii) MOLECULE TYPE: DNA (genomic)
9883 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:984:
9884 5'-CCG CGC CTC CGC CTG CCG CTT CTG
9885

*same**delete - do not show prime number.*

9886 (2) INFORMATION FOR SEQ ID NO:985:
9887 (i) SEQUENCE CHARACTERISTICS:
--> 9888 (A) LENGTH: 21 base pairs
9889 (B) TYPE: nucleic acid
9890 (C) STRANDEDNESS: single
9891 (D) TOPOLOGY: linear
9892 (ii) MOLECULE TYPE: DNA (genomic)
9893 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:985:
9894 GCT GGG CCC CGG GCG CCC CCT
9895

same

9896 (2) INFORMATION FOR SEQ ID NO:986:
9897 (i) SEQUENCE CHARACTERISTICS:
--> 9898 (A) LENGTH: 23 base pairs
9899 (B) TYPE: nucleic acid
9900 (C) STRANDEDNESS: single
9901 (D) TOPOLOGY: linear
9902 (ii) MOLECULE TYPE: DNA (genomic)
9903 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:986:

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/093,972DATE: 12/16/1999
TIME: 03:37:02

INPUT SET: S34241.raw

9904 CCC CTC TTG CTC GGG TCC CCG TG
9905*same*

9916 (2) INFORMATION FOR SEQ ID NO:988:
9917 (i) SEQUENCE CHARACTERISTICS:
--> 9918 (A) LENGTH: 23 base pairs *48 shown*
9919 (B) TYPE: nucleic acid
9920 (C) STRANDEDNESS: single
9921 (D) TOPOLOGY: linear
9922 (ii) MOLECULE TYPE: DNA (genomic)
9923 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:988:
9924 BCB GCG CGT CCT GTG TCT CCB GCB GCB TGG CCG GGC CBG CTG GGC CCC 48
9925

9946 (2) INFORMATION FOR SEQ ID NO:991:
9947 (i) SEQUENCE CHARACTERISTICS:
--> 9948 (A) LENGTH: 18 base pairs
9949 (B) TYPE: nucleic acid
9950 (C) STRANDEDNESS: single
9951 (D) TOPOLOGY: linear
9952 (ii) MOLECULE TYPE: DNA (genomic)
9953 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:991:
9954 CCC TTT TCT GGT GGG GTG
9955

18 L - more over

9986 (2) INFORMATION FOR SEQ ID NO:995:
9987 (i) SEQUENCE CHARACTERISTICS:
--> 9988 (A) LENGTH: 15 base pairs
9989 (B) TYPE: nucleic acid
9990 (C) STRANDEDNESS: single
9991 (D) TOPOLOGY: linear
9992 (ii) MOLECULE TYPE: DNA (genomic)
9993 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:995:
9994 GTG CTG TTG TTG GGC
9995

same error

SEQUENCE VERIFICATION REPORT

PATENT APPLICATION US/09/093,972

DATE: 12/16/1999
TIME: 03:37:02

INPUT SET: S34241.raw

Line	Error	Original Text
9568	Entered (23) and Calc. Seq. Length (0) differ	(A) LENGTH: 23 base pairs
9578	Entered (15) and Calc. Seq. Length (0) differ	(A) LENGTH: 15 base pairs
9588	Entered (29) and Calc. Seq. Length (0) differ	(A) LENGTH: 29 base pairs
9598	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
9608	Entered (18) and Calc. Seq. Length (0) differ	(A) LENGTH: 18 base pairs
9618	Entered (17) and Calc. Seq. Length (0) differ	(A) LENGTH: 17 base pairs
9628	Entered (15) and Calc. Seq. Length (0) differ	(A) LENGTH: 15 base pairs
9638	Entered (15) and Calc. Seq. Length (0) differ	(A) LENGTH: 15 base pairs
9648	Entered (14) and Calc. Seq. Length (0) differ	(A) LENGTH: 14 base pairs
9658	Entered (14) and Calc. Seq. Length (0) differ	(A) LENGTH: 14 base pairs
9668	Entered (18) and Calc. Seq. Length (0) differ	(A) LENGTH: 18 base pairs
9678	Entered (15) and Calc. Seq. Length (0) differ	(A) LENGTH: 15 base pairs
9688	Entered (21) and Calc. Seq. Length (0) differ	(A) LENGTH: 21 base pairs
9698	Entered (18) and Calc. Seq. Length (0) differ	(A) LENGTH: 18 base pairs
9708	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
9718	Entered (18) and Calc. Seq. Length (0) differ	(A) LENGTH: 18 base pairs
9728	Entered (21) and Calc. Seq. Length (0) differ	(A) LENGTH: 21 base pairs
9738	Entered (24) and Calc. Seq. Length (0) differ	(A) LENGTH: 24 base pairs
9748	Entered (18) and Calc. Seq. Length (0) differ	(A) LENGTH: 18 base pairs
9758	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
9768	Entered (18) and Calc. Seq. Length (0) differ	(A) LENGTH: 18 base pairs
9778	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
9788	Entered (19) and Calc. Seq. Length (0) differ	(A) LENGTH: 19 base pairs
9834	# of Sequences for line conflicts w/ running total	GGC GCC GTG CCG CGT CTT GGT GGC GGC GG
9838	Entered (30) and Calc. Seq. Length (0) differ	(A) LENGTH: 30 base pairs
9858	Entered (22) and Calc. Seq. Length (0) differ	(A) LENGTH: 22 base pairs
9868	Entered (29) and Calc. Seq. Length (0) differ	(A) LENGTH: 29 base pairs
9878	Entered (24) and Calc. Seq. Length (0) differ	(A) LENGTH: 24 base pairs
9888	Entered (21) and Calc. Seq. Length (0) differ	(A) LENGTH: 21 base pairs
9898	Entered (23) and Calc. Seq. Length (0) differ	(A) LENGTH: 23 base pairs
9918	Entered (23) and Calc. Seq. Length (48) differ	(A) LENGTH: 23 base pairs
9948	Entered (18) and Calc. Seq. Length (0) differ	(A) LENGTH: 18 base pairs
9988	Entered (15) and Calc. Seq. Length (0) differ	(A) LENGTH: 15 base pairs